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90 TTATTTTATT AATAAAATAA	180 AGCCAGTATC TCGGTCATAG	270 AATTGCATGA TTAACGTACT	360 AGTTATTAAT TCAATAATTA	450 GGCTGACCGC CCGACTGGCG	540 TGGGTGGACT ACCCACCTGA	630 AAATGGCCCG TTTACCGGGC	720 ATGGTGATGC TACCACTACG	810 GGGAGTTTGT CCCTCAAACA	900 TAGGCGTGTA CGGTGGGAGG ATCCGCACAT GCCACCCTCC
	170 CGCATAGTTA GCGTATCAAT	260 CTTGACCGAC GAACTGGCTG	350 ATTATTGACT TAATAACTGA	440 TGGCCCGCCT ACCGGGCGGA	530 TTGACGTCAA AACTGCAGTT	620 CANTGACGGT GTTACTGCCA	710 CGCTATTACC GCGATAATGG	790 800 TCCACCCCAT TGACGTCAAT AGGTGGGGTA ACTGCAGTTA	
70 CCTTTTTTT TAATTTTATT GGAAAAAAA ATTAAAATAA	160 GCTCTGATGC CGAGACTACG	250 CAAGGCAAGG GTTCCGTTCC	340 GTTGACATTG CAACTGTAAC	430 TTACGGTAAA AATGCCATIT	520 GGACTTTCCA CCTGAAAGGT	610 CTATTGACGT GATAACTGCA	700 TATTAGTCAT ATAATCAGTA		880 SAATGGGCGG TTTACCCGCC
60 GCCAGAGTAA CGGTCTCATT	150 AGTACAATCT TCATGTTAGA	240 TAAGCTACAA ATTCGATGTT	330 AGATATACGC TCTATATGCG	420 GTTACATAAC CAATGTATTG	510 ACGCCAATAG TGCGGTTATC	600 AGTACGCCCC TCATGCGGGG	690 TACATCTACG ATGTAGATGC	780 TTTCCAAGTC AAAGGTTCAG	870 CCATTGACGC GGTAACTGCG
50 GCTTCGAATA CGAAGCTTAT	140 GTCGACTCTC CAGCTGAGAG	230 GAGCAAAATT CTCGTTTTAA	320 TGTACGGGCC AGATATACGC ACATGCCCGG TCTATATGCG	410 GGAGTTCCGC CCTCAAGGCG	490 TGACGTATGT TCCCATAGTA ACTGCATACA AGGGTATCAT	590 TCATATGCCA AGTATACGGT	680 TACTTGGCAG ATGAACCGTC	770 CTCACGGGGA GAGTGCCCCT	B60 CAACTCCGCC GTTGAGGCGG
AGGCGCGCCG TCCGCGGCGCGGCGGCGGCGGCGGCGGCGGCGGGGGGGG	130 ATCCCCTATG TAGGGGATAC	220 AGTAGTGCGC TCATCACGCG	310 TGCTTCGCGA ACGAAGCGCT	400 GCCCATATAT CGGGTATATA					
30 AGGTGACCTG				390 TTAGTTCATA AATCAAGTAT	460 470 480 CCAACGACCATTG ACGTCAATAA	STO SEO STO STO STO STO STO STO STO STO STO ST	660 650 650 660 CCTGGCATTA TGCCCAGTAC ATGACCTTAT		
~ ~ ~ ~	000			380 TACGGGGTCA ATGCCCCAGT	460 470 CCGCCCATTG	STO SECULOR SECULOR STO STO STO STO STO STO STO STO STO SECULOR SECULO	CCTGGCATTA TGCCCAGTAC	GGTTTTGGCA GTACATCAAT	
30 30 30 30 30 30 30 30 30 30 30 30 30 3	110 TYTGAGATGG AGTTTGGCGC	190 200 190 200 recreecing 200 recreecing recreecing recreecing according to the second secon	290 AGAATCTGCT TAGGGTTAGG	370 370 36 36 36 370 380 380 380 380 380	460 CCAACGACCC	S50 STTTACGGTA	640 CCTGGCATTA	730 GGTTTTGGCA	820 TTTGGCACCA

Figure 14A (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement)

066	CTCTGGGTTC	1080 GCTTGCTAGC CGAACGATCG	1170 TCTGGGGGAG AGACCCCTC	1260 GTTCGCCAGA CAAGCGGTCT	1350 CGATTCACCA GCTAAGTGGT	1440 GCAAGAGGCC CGTTCTCCGG	1530 GTCTTCCCCC CAGAAGGGGG	1620 ACGGTGTCGT TGCCACAGCA	GTGGTCI	1800 GTTGGTGAGA
	TCACTATAGG (ACTGATATAGG)	1070 TCTTGCGGCC AGAACGCCGG	1160 TCTGGTGGAG AGACCACCTC	1250 CATGTATTGG GTACATAACC	1340 TGTAAAGGGT ACATTTCCCA	1430 GTATTACTGT CATAATGACA	1520 GGGCCCATCG CCCGGGTAGC	1610 CGAACCGGTG GCTTGGCCAC	1700 CCTCAGCAGC GGAGTCGTCG	1790 GGACAAGAAA . CCTGTTCTTT
	TTAATACGAC AATTATGCTG	1060 CGATTGGAAT GCTAACCTTA	1150 GTGAAGTGAA CACTTCACTT	1240 GTGACTATTA CACTGATAAT	1330 ATCCAGACAC TAGGTCTGTG	1420 ACACAGCCAT TGTGTCGGTA	1510 CTAGCACCAA GATCGTGGTT	1600 ACTACTTCCC TGATGAAGGG	1690 GACTCTACTC CTGAGATGAG	1780 ACACCAAGGT TGTGGTTCCA
096	CTTATCGAAA	1050 ACCGGTCAAT TGGCCAGTTA		1230 TYCACTTYCA AAGTGAAAGT	1320 ATAACCGACT TATTGGCTGA	1410 AAGTCTGAGG TTCAGACTCG	1500 GTCTCTGTAG CAGAGACATC	1590 CTGGTCAAGG GACCAGTTCC	1680 CAGTCCTCAG GTCAGGAGTC	1770 AAGCCCAGCA TTCGGGTCGT
950	TCCTTACTGG	1040 TCTCTAGATA AGAGATCTAT	1130 TGTTTTAAAA ACAAAATTTT	1220 AACCTCTGGA TTGGAGACCT	1310 AGGTGGTGAT TCCACCACTA	1400 GAGCCGTCTG CTCGGCAGAC	GCCAAGGGAC TCTGGTCACG CGGTTCCCTG AGACCAGTGC	1580 CCTGGGCTGC GGACCCGACG	1670 GGCTGTCCTA CCGACAGGAT	1760 CGTGAATCAC GCACTTAGTG
940	GAGAACCCAC	1030 AGGTCTCGAG TCCAGAGCTC	GCTTGGTCCT TCTTTTAAAA GGTGTCCAGT CGAACCAGA AGGAACAGGA ACAAAATTTT CCACAGGTCA	TCTCCTGTST AACCTCTGGA	1310 ACATTAGTCA AGGTGGTGAT TGTAATCAGT TCCACCACTA	1390 AACACCCTGT ACCTGCAAAT TTGTGGGACA TGGACGTTTA	1480 GCCAAGGGAC CGGTTCCCTG	1570 GCACAGGGGC CGTGTCGCCG	1660 ACACCTTCC TGTGGAAGGG	1750 ACATCTGCAA TGTAGACGTT
930		1020 ATATCTCCTT TATAGAGGAA	1110 GCTTGGTCCT CGAACCAGGA	1200 TCCCTGAAAG AGGGACTTTC	1290 TGGGTCGCAT ACCCAGCGTA	1380 AACACCCTGT TTGTGGGACA	1470 GCTTACTGGG CGAATGACCC	1560 ACCTCTGGGG TGGAGACCCC		1740 ACCCAGACCT TGGGTCTGGA
920					1270 1280 CTCCAGAGAA GAGGCTGGAG GAGGTCTCTT CTCCGACCTC	1370 CAATGCCAAG GTTACGGTTC	GGCTC	CTCCAA	1530 GGAACTCAGG CGCCCTGACC CCTTGAGTCC GCGGACTGG	
910			1100 CACCATGGAG TIGTGGTTAA GIGGTACCTC AACACCAAFT	1190 GCTTAGTGCA GCCTGGAGGG CGAATCACGT CGGACCTCCC	1270 CTCCAGAGAA GAGGTCTCTT	1360 TCTCCAGAGA	1450 TGGACGACGG GGCCTG ACCTGCTGCC CCGGAC	1540 TGGCACCCTC	1530 GGAACTCAGG	1720 TGCCCTCCAG ACGGGAGGTC
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Figure 14B (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement)

	C (1) C2	0 U U	၀ ပ ပ	9 4 H	ខ្លួក្ក	2340 IGGA ACCT	2430 IGTA ACAT	2520 CCGT GGCA	2610 GCAA CGTT	ZZ ZGAA ZCTT
	1890 AGTCCAGGGC TCAGGTCCCG	1980 TTTTCCCCAG AAAAGGGGTC	2070 GAGCCATATC CTCGGTATAG	2160 CCAGATTCCA GGTCTAAGGT	2250 CCAGGCCTCG GGTCCGGAGC	2340 GCCACATGGA CGGTGTACCT	2430 CACAGGTGTA GTGTCCACAT	2520 ACATCGCCGT TGTAGCGGCA	2610 TCTACAGCAA AGATGTCGTT	2700 ACACGCAGAA TGTGCGTCTT
	AGTC(ICAG	TTTT	GAGC	SGTO						
			2060 CCAA GGTT	2150 CCTC GGAG	2240 3CAGC	2330 CCGGA GGCCT	2420 AGAAC TCTTG	2510 CAGCG GTCGC	2600 NCTTCC AGAAGG	2690 CACAACCACT GTGTTGGTGA
	1880 ATGCAGCCCC TACGTCGGGG	1970 TCTTCTGGCT AGAAGACCGA	2060 GACCTGCCAA CTGGACGGTT	2150 TTCTCTCCTC	2240 GTAAGCCAGC CATTCGGTCG	2330 CATGTCCGGA GTACAGGCCT	2420 CCCCGAGAAC GGGGCTCTTG	2510 TATCCCAGCG ATAGGGTCGC	Z600 TCCTTCTTCC AGGAAGAAGG	CACA
				2140 CACC T					2590 CGGC GCCG	2680 TCTG AGAC
	1870 CATCCCGGCT GTAGGGCCGA	1960 AGGGAGAGGG TCCCTCTCCC	2050 GCTGGGCTCA CGACCCGAGT	2140 CTCGGACACC GAGCCTGTGG	2230 CCGTGCCCAG GGCACGGGTC	2320 TGGGTACCAA ACCCATGGTT	2410 TACAGGGCAG ATGTCCCGTC	2500 CAAAGGCTTC GTTTCCGAAG	2590 CTCCGACGGC GAGGCTGCCG	2680 TGAGGCTCTG ACTCCGAGAC
	CATO									
	1860 GACG CTGC	1950 GCTC GGAG	2040 PAGGT	2130 CTCAG	2220 GCCCA CGGGT	2310 CCACG GGTGC	2400 GTCCC	2490 CCINGCCINGET GOACGGACCA	2580 CCGTGCTGGA GGCACGACCT	2670 CCGTGATGCA GGCACTACGT
	1860 TGCCTGGACG ACGGACCTGC	1950 ACTCATGCTC TGAGTACGAG	2040 AGGGGCAGGT TCCCCGTCCA	2130 ACTCCCTCAG TGAGGGAGTC	2220 CACATGCCCA GTGTACGGGT	2310 ACACACCACG TGTGTGGTGC	2400 CCTCTGTCCC GGAGACAGGG	CCTGC	CCGT	
			2030 ACAA A IGIT I	2120 CTCC 7 3AGG 1				2480 CTGA GACT	2570 CCTC GGAG	2660 TGCT
	1830 1840 1850 CACTCC CACAGGC TCAGCGCTCC	1940 CCTCTTCACC CGGAGGCCTC TGCCCGCCCC CCACA ACTICA CGCCGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	AGGTGCCCCT AACCCAGGC CTGCACAA TCCACGGGA TTGGGTCCGG GACGTGTTT	2110 2100 2110 2110 2120 CCTAAGCCCA CCCCAAAGGC CAAACTCTCC	2210 ACAAAACTCA TGTTTTGAGT	2300 GCATCCAGGG CGTAGGTCCC	2390 GCTGTACCAA CGACATGGTT	2480 GTCAGCCTGA CAGTCGGACT	2570 ACCACGCCTC TGGTGCGGAG	2660 TYCTCATGCT AAGAGTACGA
	TCA	55	¥ 5 6	966	3 PC 0 0					
	1840 2AGGC 3TCCG	1930 300TC	2020 AGGCC TCCGG	2110 AAGGC TTCCG	2200 TTGTG	2290 PAGCCT	2380 3TGACC	2470 CAAGAACCAG GITCTIGGIC	2560 CAACTACAAG GTTGATGTTC	2650 GGGGAACGTC CCCCTTGCAG
-		0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	AACCCAGGCC TTGGGTCCGG	2000 9000 1000	2200 2190 AAATCTTGTG	CAGGIGGE AGAGIAGET CAGGIGA DETCATEGGA	2380 GAGAGTGACC CTCTCACTGG	CAAG		
	1830 RECTO O	1920 CACC O	2010 - 20CCT 7	2100 CCCA GGGT	2190	2280 CCCT		2460 TGAC	2550 AGAA TCTT	2640 AGCA
٠	1830 GTGTCTGCTG	1920 CCTCTTCACC	2010 AGGTGCCCT TCCACGGGGA	AAGC TTCG	TGCAGAGCCC	CAGGTGCCT CAGGTGCCCT	2370 CCTCTGCCCT	2460 ATGAGCTGAC TACTCGACTG	2550 AGCCGGAGAA TCGGCCTCTT	2640 GGTGGCAGCA CCACCGTCGT
	1820 GAGG	1910 1910 1910 1910 1910 1910	2000 AGGCT	2090 2090 2090	2180 CTCTC	2270 2270 2270 2000A	. 2360 3CCCAC	2450 CCATCCCGGG	2540 ATGGGC	2630 GACAAGAGCA CTGTTCTCGT
•	GGAGG		GCAC		ATCTT	2270 CAAGGCGGGA	CTCGGCCCAC		AGCAAT	
	1810 GGCCACA GGGAGGGAGG	ACCANGECAG GCCCCGTCTG	CCTCTGGGCA GGCACAGGCT	CGGGAGGACC CTGCCCTGA	2170 2180 CTARCTCTCTC CTARGET				GCAGTGGGAG ACCAATGGGC	2620 2620 2630 CCTCACCGTG GACAAGAGACA CGAGTGGCAC CTGTTCTCGT
	CAGC	AAGG		GGAGG	AACTO		2350 2350 CAGAGGCCGG	GTCTCCGGCC 2440 CACCCTGCCC	GAGTO	CTCAC
	9 8	3		3 88	3 8	5 8	5 D	5 01	. 00	ט פי כ

Figure 14C (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement)

TGCTTGGCAC ACGAACCGTG	2880 ATGGTTCTTT TACCAAGAAA	2970 TGTGCAGGTG ACACGTCCAC	3060 AGCAGCACCT TCGTCGTGGA	3150 TTCTGTGAGC AAGACACTCG	3240 CTACCCCCAC GATGGGGGTG	3330 CCTGTGGAGG GGACACCTCC	3420 CACCACACAC GTGGTGTGTG	3510 GAACACTCCT CTTGTGAGGA	3600 TCAGACAAAC AGTCTGTTTG	
TGCTT	ATGGI	TGTG	AGCA	TTCT				_		
2780 CGCACGAGGA GCGTGCTCCT	2870 CGAGACTGTG GCTCTGACAC	2960 TGGCCCAGGC ACCGGGTCCG	3050 CCCTCCCTCC GGGAGGGAGG	3140 GACTGTCCTG CTGACAGGAC	3230 CCTCACCCAT GGAGTGGGTA	3320 ACTCTCGGGC TGAGAGCCCG	3410 GCCACACGGC CGGTGTGCCG	3500 TCGCACACGT AGCGTGTGCA	3590 GCTGACCTGC CGACTGGACG	
2770 CTCTCGCGGT (GAGAGCGCCA (2860 TGGGCCCCTG ACCCGGGGAC	2950 GTCCCCACAC CAGGGGTGTG	3040 GCCAGCGTGG CGGTCGCACC	3130 CTCTGTAGGA GAGACATCCT	3220 ACAGGCCCTC TOTCCGGGAG	3310 GGGGACATGC CCCCTGTACG	3400 AGGTTGGCCG TCCAACCGGC	3490 AGCAAGGTCC TCGTTCCAGG	3580 TTCTCCACAT AAGAGGTGTA	
2760 GCTCCCCGGG CGAGGGGCCC			3030 TGGGGGATTT ACCCCCTAAA	3120 CAGCCCCTGC GTCGGGGACG	3210 GTGCGTAGGG CACGCATCCC	3300 AACCGACTCC TTGGCTGAGG	3390 CCCCGCACTG GGGGCGTGAC	3480 CCCAGACCAG GGGTCTGGTC	3570 TCTCGGCAGC	
	2840 TAAAGCACCC ATTTCGTGGG	2930 GAGGCAGAGC CTCCGTCTCG	3020 CTCGGCAGGG GAGCCGTCCC	3110 GACAGACACA CTGTCTGTGT	3200 CCTAGICCAT GGAICAGGIA	3290 ATGGGGACAC TACCCCTGTG	3380 GTTCAACAAA CAAGTTGTTT	3470 CTGCACAGCA GACGTGTCGT	3560 CCCACGAGGC GGGTGCTCGG	
GTAAATGAGT GCGACGGCCG GCAAGCCCCC	2820 2820 2830 2840 2850 CCGGGGCGCCC AGGATGGAAA TAAAGCACCC AGCGTACCTTT ATTTCGTGGG TCGCGACGG	2920 TGGCATGAGG ACCGTACTCC	3010 AGGGCTGCC TCCCCACGG	3100 AGCCCCTGGG TCGGGGACCC	3190 CGGGGGCATG GCCCCGTAC	3280 TCGCACCCGC AGCGTGGGCG	3370 GCCCAGACCC CGGGTCTGGG	3460 CCCGGGGGAA GGGCCCGATT		
GTAAATGAGT GCGACGGCCG	2820 2830 CCGGGCGCC AGGATGGAAA GGCCCGCGG TCGTACCTTT	2910 GAGGCCTGAG CTCCGGACTC	3000 GGGCTCAGCC CCCGAGTCGG	3090 AAGCCCTAGG TTCGGGATCC	3180 CATGCCCACT GTACGGGTGA	3270 CTGCCCAGCC GACGGGTCGG				•
			2990 CCCTAGGGTG GGGATCCCAC	3080 GGGCCACGGG	3170 TCCCGACCTC	3260 CCTGGCTGCC GGACCGACGG	3350 GATGCCCACA CTACGGGTGT			
GAGCCTCTCC CTGTCTCCGG	2810 2800 2810 GTACCCCCTG TACATACTTC CATGGGGGAC ATGTATGAAG	2890 CCACGGGTCA GGTGCCCAGT	2980 TGCCTGGGCC	3070 GCCCTGGGCT	3160 GCCCTGTCC GCCCTGTCC	3250 GGCACTAACC	3340 GACTGGTGCA GATGCC	3430 ACACGTGCAC	3520 CGGACACAGG GCCTGTGTCC	
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Figure 14D (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement)

3690 TGGCCCACTT ACCGGGTGAA	3780 CCCGTGCCTT GGGCACGGAA	3870 CATTCTATTC GTAAGATAAG	3960 ATGGCTTCTG TACCGAAGAC	4050 GTTACGCGCA CAATGCGCGT	4140 CCTCTCAAAA GGAGAGTTTT	4230 CCCAGTTCCG GGGTCAAGGC	4320 AAGTAGTGAG TYCATCACTC	4410 TCCTAGCGTG AGGATCGCAC	4500 ATTGGCAAGA TAACCGTTCT
3680 TCCCTGGCCC 7	3770 TTGCCCCTCC	3860 GAGTAGGTGT CTCATCCACA	3950 GGTGGGCTCT CCACCCGAGA	4040 GGGTGTGGTG CCCACACCAC	4130 GTTCGCCGGG CAAGCGGCCC	4220 CCTAACTCCG GGATTGAGGC	4310 GCTATTCCAG CGATAAGGTC	4400 TYGACGGCAA AACTGCCGTT	4490 AAATATGGGG TTTATACCCC
3670 CCACGTCACG GGTGCAGTGC	3760 CATCTGTTGT GTAGACAACA	3850 CGCATTGTCT GCGTAACAGA	3940 CTGGGGATGC GACCCCTACG	4030 TAAGCGCGGC ATTCGCGCCG	4120 TTCTCGCCAC AAGAGCGGTG	4210 CCATCCCGCC GGTAGGGCGG	4300 CGGCCTCTGA GCCGGAGACT	4390 CGCGCCAAAC GCGCGGTTTG	4480 CCGTGTCCCA GGCACAGGGT
3660 GGATCACACA CCTAGTGTGT	3750 AGTTGCCAGC TCAACGGTCG	3840 GAAATTGCAT CTTTAACGTA	3930 AGCAGGCATG TCGTCCGTAC	4020 AGCGGCGCAT TCGCCGCGTA	4110 TTCCCTTCCT AAGGGAAGGA	4200 CTAACTCCGC GATTGAGGCG	4290 GAGGCCGCCT CTCCGGCGGA	4380 GCTCCGATTT CGACGCTAAA	4470 TGCATCGTCG ACGTAGCAGC
3650 CACACACAGG GTGTGTGTCC	3740 TGTGCCTTCT ACACGGAAGA	3830 ATAAAATGAG TATTTTACTC	3920 GGAAGACAAT CCTTCTGTTA	4010 CGCGCCCTGT GCGCGGGACA	4100 TTTCGCTTTC AAAGCGAAAG	4190 AGTCCCGCCC TCAGGGCGGG	4280 TGCAGAGGCC ACGTCTCCGG	4370 ACAGCTCAGG TGTCGAGTCC	4460 ACCATTGAAC TGGTAACTTG
	3730 CAGCCTCGAC GTCGGAGCTG	3820 TCCITTCCTA AGGAAAGGAT	3910 GGGAGGATTG CCCTCCTAAC	4000 GGTATCCCCA CCATAGGGGT	AGCGCCCTAG CGCCCGGTCC TCGCGGATC GGGGGGGAAGG	4180 CAGCAACCAT GTCGTTGGTA	4270 TTTTATTTA AAAAATAAAT	4360 AAAAGCITGG TYTTCGAACC	4450 TCATGGTTCG AGTACCAAGC
orecected accected cacegearer	CAGGACGGAT CAGCCTCGAC GTCCTGCCTA GTCGGAGCTG	3810 11 3820 ACTCCCACTG TCCTTTCCTA TGAGGGTGAC AGGAAAGGAT	3900 GACAGCAAGG CTGTCGTTCC	3990 GGCTCTAGGG CCGAGATCCC	4080 AGCGCCCTAG TCGCGGGATC	4170 CTCAATTAGT GAGTTAATCA	4260 TGACTAATTT ACTGATTAAA	4350 GGCTTTTGCA CCGAAAACGT	4430 GGALTITAIC CCCGCIGCCA CCIAAAAIAG GGGCGACGGI
	3710 CCCTTCCCTG GGGAAGGGAC		3890 GGTGGGGCAG CCACCCGTC	3980 AACCAGCTGG TTGGTCGACC	4070 GCGTGACCGC TACACTTGCC CGCACTGGCG ATGTGAACGG	4150 AAGGGAAAA AAGCATGCAT TTCCCTTTTT TTCGTACGTA	4250 GCCCCATGGC CGGGGTACCG	4330 4340 4350 GAGGCTTYTT TGGAGGCCTA GGCTTTTYGCA CTCCGAAAAA ACCTCCGGAT CCGAAAACGT	AAGGCTGGTA GGATTTTATC CCCGCTGCCA TTCCGACCAT CCTAAAATAG GGGCGACGGT
3620 CCAGCCCTCC TCTCACAAGG GGTCGGAAGG AGAGTGTTCC	3700 CCCAGTGCCG GGGTCACGGC	3790 3800 CCTTGACCCT GGAAGGTGCC GGAACTGGGA CCTTCCACGG	1880 10GGGGGTGG GGTGGGGCAG ACCCCCCAC CCACCCGTC	3970 AGGCGGAAAG TCCGCCTYTC	4060 GCGTGACCGC TACACT CGCACTGGCG ATGTGA	4150 AAGGGAAAAA AAGCAT TTCCCTTTTT :TCGTA	4240 CCCATTCTCC GGGTAAGAGG	4330 GAGGCTTTTT CTCCGAAAAA	4420 AAGGCTGGTA TTCCGACCAT
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Figure 14E (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement)

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4680 AGTAGAGAAC ICATCTCTTG	4770 GCAAGTAAAG CGTTCATTTC	4860 ACAAGGATCA TGTTCCTAGT	4950 CTCTCTGAGG GAGAGACTCC	5040 GCTCCCCTCC CGAGGGGAGG	5130 TGACATAATT ACTGTATTAA	5220 TAATYGTTYG ATTAACAAAC	CAGAAG	5400 AGGACTTTCC TCCTGAAAGG
		4850 ACTCTTTGTG TGAGAAACAC	4940 CCCAGGCGTC GGGTCCGCAG	5030 CAAGTTCTCT GTTCAAGAGA	5120 TTCTGTGGTG AAGACACCAC	5210 CTACTGATTC GATGACTAAG	_	5390 GAAGACCCCA CTTCTGGGGT
	4750 TTATTGAACA AATAACTTGT	4840 GCCACCTTAG CGGTGGAATC	4930 TCCCAGAATA AGGGTCTTAT	5020 AAGATGCTTT TTCTACGAAA	5110 GGAACCTTAC CCTTGGAATG	5200 ATGTGTTAAA TACACAATTT		5380 GAGAAAGGTA CTCTTTCCAT
		4830 AATCAACCAG TYAGTTGGTC	4920 TATAAACTTC ATATTTGAAG	5010 GACTAACAGG CTGATTGTCC	5100 TCTTTGTGAA AGAAACACTT	5190 TAAGTGTATA ATTCACATAT		5370 Caaaaaagaa Gititititciti
		4820 GGAAGCCATG CCTTCGGTAC	4910 TTTGGGGAAA AAACCCCTTT	5000 CGAGAAGAAA GCTCTTCTTT	5090 GCTTTAGATC CGAAATCTAG			5360 TCTACTCCTC AGATGAGGAG
4630 CCATTCCTGA GGTAAGGACT	4720 THECCAAAAG	4810 CTGTTTACCA GACAAATGGT	CAGAAATIGA GICTTIAACT	4990 TITGAAGTCTA AACTTCAGAT	5080 ACTITITGCTG TGAAAACGAC			5350 CTCTCAACAT GAGAGTTGTA
		4800 GGAGGCAGTT CCTCCGTCAA	4890 ACGITITICC. TGCAAAAGG	4980 AAGTATAAGT TTCATATTCA,	5070 AGACCATGGG TCTGGTACCC	5160 TTTAAAGCTC AAATTTTCGAG		5340 CTACTGCTGA GATGACGACT
		4790 TYGGATAGTC AACCTATCAG	4880 TGAAAGTGAC ACTTTCACTG	4970 AAAAGGCATC TTTTCGTAG			5240 ATTCCAACCT TAAGGTTGGA	5320 GCCATCTAGT GATGATGAGG CGGTAGATCA CTACTACTCC
4600 TCGTGATTAT (4690 TCAAAGAACC AGTTTCTTGG	4780 TAGACATGGT ATCTGTACCA	4870 TGCAGGAATT ACGTCCTTAA	4960 TCCAGGAGGA AGGTCCTCCT	5050 TAAAGCTATG	5140 GGACAAACTA CCTGTTTGAT	5230 TGTATTTTAG ACATAAAATC	5320 GCCATCTAGT CGGTAGATCA
	GGGTAGGAAA ACCTGGTTCT CCATTCCTGA GAAGAATCGA CCTTTAAAGG ACAGAATTAA TATAGTTCTC ACTTCTC CCCATCCTTT TGGACCAAGA GGTAAGGACT CTTCTTAGATTCC TGTCTTAATT ATATCAAGA TCATCTT	ACCTGGTTCT CCATTCCTGA GAAGAATCGA CCTTTAAAGG ACAGAATTAA TATAGTTCTC ACTAGACTGACCAAGA GATAAGGACT CTTCTTAGCT GGAAATTTCC TGTCTTAATTT ATATCAAGAG TCATCTCTCTCTCTCTCTCTCTCTCTCTCTTAATTT ATATCAAGAG TCATCTCTCTCTCTCTCTCTCTCTTAATTTCTTCTTAATTGATTG	ACCTGGTTCT CCATTCCTGA GAAGAATCGA CCTTTAAAGG ACAGAATTAA TATAGTTCTC AGTAGAG TGGACCAAGA GOTTCCTGA GAAGAATCGA CCTTTAAAGG ACAGAATTAA TATAGTTCTC AGTAGAG TGGACCAAGA GOTTTGGATGAT GGAAATTCTC TGTCTTAATT ATATCAAGAG TCATCTC 4710 4710 4710 4710 4710 4710 4710 471	ACCTGGTTCT CCATTCCTGA GAAGAATCGA CCTTTAAAGG ACAGAATTAA TATAGTTCTC AGTAGAG TGGACCAAGA GATAO TTTCGAATTCC TGTCTTAATT ATATCAAGAG TCTCTTAGATTCC TGTCTTAATT ATATCAAGAG TCATCTC GGAATTCC TGTCTTAATT TCTGAAAAG TTTCGAAAAG TTTCGAATTCTA AGGATTCTG AGAGAATTCTG ACGGAATTG GCAAGAA GGAGGCCATG AABACCTACTA CGGAATTCTG AAAACTTCT TGGCCAAAAG TTTGGGGAA TCTTCGGTAC TTAGTTGGTC CGTGGAATC TGAGAAAACA TTTGGGGAAA TAGTTGGGAAA TAGTTGGGAAA TAGTTGGGAAAT TTTGGGGAAA TAGTTGGGAAATCC TGAGAAAACA TTTGGGGAAA TAGTTGAACTTC TCCCAGAATCC TGAGAAAACA TTTGGGGAAA TATAGAAATTGA TTTGGGGAAA TATAGAAATTGA TTTGGGGAAA TATAGAAATTGA TTTGGGGAAA TATAGAAATTGA TTTGGGGAAA TGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTC	ACCTEGATICE CONTECTED GAAGAATICA GAAGAATITAC TATACATICA ACTOR GAAGAATICA GAAAATITAC TACCAAAAA TATACATACA ACCGGAATITAC ACTACAAAAA TATACAAAAA TATACAAAAAA TATACAAAAAA TATACAAAAAAAA	ACCTEGITCT CCATTCCTGA GAAGAATCGA CCTTTAAAGG ACAGAATTAA TATACTTCTC A TGGACCAAGA TTCCAAGAG TTCCTGA GAAAATTCC TGCATTCCTGA GAAGAATCGA GAAAATTCC TGCTTAAATT ATATCCAGGA TTCCAAAAG TTCCAAAAG TTCCAAAAG TTCCAAAAG TTTCCAAAAG TTTCCAAAAG TTTCCAAAAGA TTTCAACCAGGACTTC AACCTTCAACA GACCATTAACCA GAAACCTACTC GAAATTCAA GACCTTTAACCA GAAAACCTAC TTACTCAACA GACCATTAACCA GAAAATTCAA TTTCCAAAAACG TTTCAAAAAGG GCCACCTTTAA GACCTTTAACT TTCCAAAATTCA TTTCCAAAAACG TTTCAAAAACG GCCACCTTTAA GACCTTTAACT TTCCAAAAAAGG GCCACCTTTA TTTCAAAAAAAGG GACCATTTTCAAAAAAAAAA	AGENCIATION 4650 4660 4670 ACCTICALTITIC CAPTICCTCA GARGATITICC TGTCTTAATT ATATICANGG TTGCACCATCA GARGATITICC TGTCTTAATT ATATICANGG TTGCACCATCA GARGATITICC TGTCTTAATT ATATICANGG TTGCACCATCA GARGATITICC TGTCTTAATT ATATICANGG TTGCACCATCA GARGATITICC TGTCTTTAATT ATATICANGG TGTCTTTAC GGARTTCTG ATATICANGG TTTGCACCATCA GGARTTCTG ATATICANGG TTTGCACCATCA GGARTTCTG ATATICANGG TTTGCACCATCG TGACCTTTAC GGARTTCTG TGGCCTTAAC GGARTTCTG TGGCCTTAAC GGARTTCTG TGGCCTTAAC GGARTTCTC TGGCTTTTACC GGARTTCTC TTGCACAAATGG TTTGGGGAAA TATGGGGAAA TATGGGGAAA TATGGGGAAA TATGGGGAAA TATGGGGAAA TATGAGGAAA TATGGGGAAA TATGGGGAAA TATGGGGAAA TATGGGGAAA TATGGGGAAA TATGGGGAAA TATGAGGAAA TGGTTAAACTCT TGCACAAATGG TTGAAAATGG TTGAAAAAGG GTCTTTTAAACTT TGCAAAAAGG GTCTTTTTTAAACT TTGAAATTCTT GGTCCGCAG TGCAAAATGGT TTGAAAAAGG GTCTTTTTTTTTT	ACCUTAGATOR GARAGAATOR CCTTTAAAGG ACAGAATTAA TATACAAGAG TAGACCCAAGA GATAAGACCCAAGA GATAAGACCCAAGA GATAAGACCCAAGA GATAAGACCCTTAAGA GATAAGATTAC GAAATTTCC TGTTTATAAAG TTTGGAAATTAC GGAAATTTCC ATTAGAAAG AAGCCTTAAGA GATAACCTTAACA GAAGCCATTAC GGAAATTCCA GAAGCCATTAC GGAAATTCCA GAAGCCATTACA GAAGCCATTACA GAAGCCATTACA GAAGCCATTA CCTTCAAATGGT CCTCCGTTAAAAG GACCATTAAAAATTCCA GGAAGCCATTA ATTAGAGAATTCCA GAAGCCATTA ATTAGAGAAAAAAAAAA

Figure 14F (SEQ ID NO.: 10 - Primary Sequence) (SEQ ID NO.: 28 - Complement)

5490 AAAAAGCTGC TTTTTCGACG	5580 TTTTTCTTAC AAAAAGAATG	5670 TTAATAAGGA AATTATTCCT	5760 CTCCCACACC GAGGGTGTGG	5850 AGCAATAGCA TCGTTATCGT	5940 GTCTGGATCG CAGACCTAGC	6030 TACAAATAAA ATGTTTATTT	6120 TCTTATCATG AGAATAGTAC	6210 ACAATTCCAC TGTTAAGGTG	5300 TCACTGCCCC AGTGACGGG(
5480 ACCACAAAGG AA TGGTGTTTCC TT	5570 AACATACTGT TT TTGTATGACA AA	5660 TGTAAAGGGG TT ACATTTCCCC AA	5750 TITIAAAAAAC C. AAATITITITIG GI	5840 TTACAAATAA M AATGTTTATT T	5930 ATCTTATCAT G TAGAATAGTA C	6020 TTATAATGGT T AATATTACCA	6110 CATCAATGTA TCTTATCATG GTAGTTACAT AGAATAGTAC		6290 TGCGTTGCGC ACGCAACGCG
5470 TGCTATTTAC A ACGATAAATG T	5560 TTATAATCAT A AATATTAGTA T	5650 CTTTTTAATT 1 GAAAAATTAA 1	5740 TTTTACTTGC	5830 CTTATAATGG GAATATTACC	5920 TCATCAATGT AGTAGTTACA	CCCAACTYGT TTATYGCAGC GGGTYGAACA AATAACGTCG	AGTIGIGGTT TGTCCAAACT TCAACACAA ACAGGTTTGA	6190 TGTGAAATTG ACACTTTAAC	
5460 TTGCTTGCTT 1		5640 GTACCTTTAG CATGGAAATC	5730 TYTGTAGAGG AAACATCTCC	5820 TTTATTGCAG AAATAACGTC	5910 TYGTCCAAAC AACAGGTTYG		6090 AGTTGTGGTT TCAACACCAA	6180 CTGTTTCCTG GACAAAGGAC	6270 GTGAGCTAAC CACTCGATTG
			5720 CCATACCACA GGTATGGTGT	5810 TGTTAACTTG ACAATTGAAC	5900 TAGTTGTGGT ATCAACACCA			6170 ATGGTCATAG TACCAGTATC	6260 TGCCTAATGA ACGGATTACT
5450 TCTGTTTAGT AATAGAACTC	5530 Trongtaacc	5620 5620 TAACTATGT CAAAAATTGT ATTGATACGA GITTITTAACA	and the same of th	5800 CAATTGTTGT GTTAACAACA	S890 CACTGCATTC GTGACGTAAG	5980 TGCTGGAGTT ACGACCTCAA	-		6250 AAGCCTGGGG
5430 5440 5430 5450 5450 5450 5450 5450	S520 5540 5540 TOGAAAAATA TAGAAATTGG AAATATTCAT	5610 5620 5620 CTGCTATTAA TAACTATGCT GACGATAATT ATTGATACGA	5700 TGACTAGAGA	5790 AAAATGAATG	S880 GCATTTTTT CGTABABABA	5970 GGGGATCTCA			
			5690 GCCT		SEGUESACTI CONCILIERA 5860 5870 TCACAAATT CACAAATAAA	GCTGGATGAT CCTCCAGCGC	CGACCIACIA GGAGGICGCG 6040 GCAATAGCAT CACAAATTTC COMMANGGATA GREATTTAAAA	COLLEGIA CONTROL 6140 TCTGTATACC GTCGACCTCT AGACATATGG CAGCTGGAGA.	6230 ACAACATACG AGCCGGAAGC TGTTGTATGC TCGGCCTTCG
5410 TTCAGAATTG	AAGICITAAC GAIICAAAAA 5500 S510 ACTGCTATAC AAGAAAATTA	SSSO SSSO SSSO TCCACACAGG CATAGAGTGT AGGINGTICTCACACACAGG CATAGAGTGTCACACACACACACACACACACACACACACA	S680 S690 ATATTIGATIC TATAGEGGA	TCCCCTGAA CCTGAAACAT	5860 TCACAAATTT	S950 GCTGGATGAT	CGACCIACIA 6040 GCAATAGCAT	6130 TCTGTATACC	6220 ACAACATACG TGTTGTATGC
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Figure 14G (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement)

6390 CGCTCTTCCG GCGAGAAGGC	6480 TATCCACAG ATAGGTGTC	6570 CGTTTTTCC GCAAAAAGG	6660 TACCAGGCGT ATGGTCCGCA	6750 GGAAGCGTGG CCTTCGCACC	6840 CCCGTTCAGC GGGCAAGTCG	6930 ACTGGTAACA TGACCATTGT	7020 GTATTTGGTA CATAAACCAT	7110 GGTGGTTTTT CCACCAAAAA	AGACGCAGA CGACTICGGT CARIOGRAD 7150 7160 7170 7180 7200 7200 7200 7200 7200 7200 7200 72
6380 CGTATTGGG CC GCATAACCC G	6470 IGTAATACGG T CATTATGCC A	6550 6550 GCGTTGCTG GCGTTTTTCC GTAAAAGGC CGCGTAACGAC CGCAAAAAAGGC CGCAAAAAAGG	6630 6640 6650 6650 ACCCAGAGA ACTATANAGA TACCAGGGGT AGGTGGCGAA ACCCGACAGG ACTATANAGA TACCAGGCGT AGGTGTCC TGATATTTCT ATGGTCCGCA	CTCGTAGTGT TTTTAGCTGC GAGTICAGIC ICCACCOCT 6730 6730 6740 6750 6750 6750 6750 6750 6750 6750 675	AAGGGGACC TYCGAGGGAG CACGCGAAGAG GACAACGCGAAA 18800 6810 6820 6830 6830 6840 6770 6780 6770 6780 GYAGTYCGGY GYAGGYCGYY CACTCCAAGC TGGCCYGYGY CCACGAAGYCG GGCAAAGYCG	GCGAAAGAGT TACGAGTGCG ACATCCATAG AGTCAAGCCA CAICCAGCAAAGAGT TACGCCACT GGCAGCAGCA 6910 6950 6950 6950 6950 6950 6950 6950 695	GGCTGGCGAC GCGGAATAGG CCATTGATAG CAGAACTCAG G11555CAT 1050 7000 7010 7020 7020 6940 6950 6950 6950 6950 GTATTTGGTA GGATTAGCTA ACGCCTACAC TAGAAGACA GTATTTGGTA GGATTAGCAG AGCGAAGAACTAG ACCGAATGA GTAGGCGGTG CTACAAGAAGAG ACCGAATAGCAG AGCGAAGAGA TGCCGAATGA ACCCAATGAA ACCGAATAGAAGAAGAAGAAGAAGAAGAAAACAATAGAAAAAAAA	CCTAATCGTC TCGCTCCATA CATCGCCAC GALGICICAN OFFICE TOTO 7080 7090 7090 7100 71100 71100 71100 71100 71100 71100 71100 7100 7030 703	7190 GTCTGACGCT CAGACTGCGA
6370 REGCEGITT G	6460 ACTCAAAGGC G	6550 GTAAAAAGGC CATTTTTCCG	6640 ACCCGACAGG TGGGCTGTCC	6730 TOTCCGCCTT	6820 TGGGCTGTGT ACCCGACACA	. 6910 TATCGCCACT ATAGCGGTGA	7000 ACGCCTACAC TGCCGATGTG	7090 A AACAAACCAC F TTGTTTGGTG	7180 F TTTCTACGGG A AAGATGCCC
6369 GCGCGGGGA DCCGCCT	6450 TRATCAGCTC P	6540 GCAGGAACC	6630 AGGTGGCGAA	6720 ACCGGATACC	6810 CGCTCCAAGC	6900 AGACACGACT	6990 TGGCCTAACT ACCGGATTGA	1080 TGATCCGGCA	1170 r cctttgatct a ggaaactaga
6350 ATCGGCCAA C	6440 CGGCGAGCG	6530 CAGCAAAAG	6620 CTCAAGTCAG	GAGTICAGIO 6710 CCTGCCGCTT	GGACGGCGAA 6800 GTAGGTCGTT	GTCAAGCCA CAICCAGCAG 6880 6880 TCTTGAGTC CAACCGGGTA	6711646CCCA 6980 CTTGAAGTGG	TGGTAGCTCT	TCAAGAAGAT AGTTCTTCTA
6340 GCATTAATG A	6430 OCTTCGGCT	GCAAGCCGA 6520 3AGCAAAAGG	STCGTTTTCQ 6610 AAAATCGACG	TTTTAGCTGC 6700 CTGTTCCGAC	GACAAGGCTG 6790 TCAGTTCGGT	AGTCAAGCCA 6880 GTCTTGAGTC	CAGAACTCAG 6970 CTACAGAGTT	7060 GAAAAAGAGT	7150 AAAAAGGATG TTTTTCCTAG
6330 T DECCAGG	GCACGGTCG A 6420 TGCGCTCGG T	ACGCGAGCC P 6510 AGAACATGT (TCTTGTACA 6600	CTCGTAGTGT 6690 GTGCGCTCTC	CACGCGAGAG 67B0 TGTAGGTATC	ACATCCATAG 6870 GGTAACTATC	CCATTGATAG 6960 GTAGGCGGTG	CCTAATCGTC TCGCTCCATA CATCCGCCACACACACACACACACACACACACACACACAC	AGACGCGAGA CGACTICGGI CANTGGARGO 7120 7130 7140 TIGITIGGAA GCAGCAGATT ACGCGCAGAA AACAAACGTT CGTCGTCTAA TGCGCGTCTT
6320 GGAAACCTG T	CCTTTGGAC A 6410 ACTGACTCG C	TCACTGAGC G 6500 PACGCAGGA A	MTGCGTCCT 1 6590	SGGGGGACTG 6680	AAGGGGACC TTCGAGGGAG 6760 6770 CGCTTTCTCA ATGCTCACGC	TACGAGTGCG 6860 CGCCTTATCC	GCGGAATAGG 6950 AGCGAGGTAT	TCGCTCCATA 7040 GCTGAAGCCA	CGACTICGGT 1130 CGCAGCAGATT CGTCGTCTAA
6310 6320 6330 6340 6350 6350 6350 6310 6320 6320 6320 6320 6320 6320 6330 633	GAAAGGTCAG CCCTTTGGAC AGCACGGTCG ACGTAATTAG 11A0CUGGT 6440 6450 6450 6460 6460 6410 6410 6450 6400 6410 6410 6410 6450 6410 6410 6410 6410 6410 6410 6410 641	GAAGGAGCGA GTGACTGAGC GACGCGAGCC AGCAAGCCGAAGCG 6530 6530 6540 6500 6500 6500 6500 AATCAGCAAGAA CCCAGGAAAGG CCAGGAAAAG GCCAGGAAACCC AATCAGGGAA TAACGCAGGA AAGAACATGT GAGCAAAAGG CCAGCAAAAG GCCAGGAAACC	THAUTCCCCT ATTGCGTCCT TTCTTGTACA CTCGTTTTCG GGICLOSTCCCT 6630 6640 6650 6650 6650 6650 6650 6650 665	TATCCGAGGC GGGGGACTG CTCGTAGTGT TTTTAGCTGC GAGTICAGIC ICACCGGGATACC 6730 6740 6720 6730 6740 6770 6770 6770 6770 6770 6770 677	AAGGGGGACC 6760	GCGAAAGAGT 6850	GGCTGGCGAC 6940 GGATTAGCAG	CCTAATCGTC 7030 TCTGCGCTCT	AGACGCGAGP 7120 TYGTTYGCAJ

Figure 14H (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement)

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rttaaatca Vaatittagt	7380 TTTCGTTCAT AAAGCAAGTA	7470 CCGCGAGACC GGCGCTCTGG	7560 TCCGCCTCCA AGGCGGAGGT	7650 ACAGGCATCO TGTCCGTAGC	TTGTGC	CTGCATI	CGGCGAC	8010 TCTTCGGGGC	B100 TTTACTTTCA
7250 ACCTAGATCC TTTTAAATTA AAANTGAAGT TTTAAATCAA TGGATCTAGG AAAATTTAAT TTTTACTTCA AAATTTAGTT	7370 GATCTGTCTA '	7460 TGCAATGATA ACGTTACTAT	7560 7550 7550 7560 GAAGTGGTCC TGCAACTTTA TCCGCCTCCA CTTCACAG ACGTTGAAAT AGGCGGAGGT	7620 7630 7630 7630 ACAGGCATCG ACAGGCATCG TIATCAAACG CGTTGCAACA ACGGTAACGA TGTCCGTAGC	7720 7730 GAGTTACATG ATCCCCCATG CTCAATGTAC TAGGGGGTAC	7820 TATGGCAGCA ATACCGTCGT	7910 7900 7900 TCAACCAAGT CATTCTGAGA ATAGTGTATCA GTAAGACTCT TATCACATAC	TGCTCATCAT TGGAAAACGT ACGAGTAGTA ACCTTTTGCA	8090 TTCAGCATCT
7270 TITTRAATTTA AAAATTTAAT	AGTOAGGCAC CTATCTCAGC GATCTGTCTA TCACTCGTG GATAGAGTCG CTAGACAGAT	7450 GCCCCAGTGC CGGGGTCACG	7540 GAAGTGGTCC CTTCACCAGG	7630 GCAACGTTGT CGTTGCAACA		GCAGTGTTAT CACTCATGGT CGTCACAATA GTQAGTACCA	7900 CATTCTGAGA GTAAGACTCT	1990 TGCTCATCAT:	BOBO CCAACTGATC GGTTGACTAG
ACCTAGATCC TGGATCTAGG	7350 AGTGAGGCAC TCACTCCGTG	7440 TTACCATCTG AATGGTAGAC	7530 GCCGAGCGCA CGGCTCGCGT	7620 AATAGTTTGC TTATCAAACG	7710 CGATCAAGGC GCTAGTTCCG			7980 ACTTTAAAAG TGAAATTTTC	8070 ACTCGTGCAC TGAGCACGTG
	7340 ATGCTTAATC FACGAATTAG	7430 ACGGAGGGC TGCCCTCCCG	7520 AGCCGGAAGG TCGGCCTTCC		CGGTTCCCAA GCCAAGGGTT	TICTCACADA PARGITGGCC AACAGICTIC ATCAACCGG	7880 TGGTGAGTAC	ATACCGCGCC ACATAGCAGA TATGGCGCGG TGTATCGTCT	B060 GATGTAACCC CTACATTGGG
7240 7250 7250 GATTATCAAA AAGGATCTTC CTAATAGTTT TTCCTAGAAG	7310 7310 7310 7320 7330 7340 7340 7340 7340 7340 7340 734	7420 7420 7450 7450 TAACTACCATCTG GCCCCAGTGC TGCAATGATA ATTGATGCTAGTGC AATGGTAGAC CGGGGTCACG ACGTTACTAT	CACGCTCCACGAT TTATCAGCAA TAAAACCAGC AGCCGGAAGGGGGGGAGTGG CGAGGTCTA AATAGTCGTT ATTTGGTCGG TCGGCTTCC	7600 GAGTAAGTAG CTCATTCATC	CATTCAGGTC GGGTTCCCAA GTAAGTCGAG GCCAAGGGTT		7880 7860 7880 7880 GTAAGATGCT TTTCTGTGAC; TGGTGAGTAC CATTCTACGA AAAGACACTG ACCACTCATG	7930 7940 7950 7950 7950 7950 7970 7970 7980 7980 8010 8010 8010 8010 8010 8010 8010 8	8020 8030 8040 8050 8050 8060 8060 8070 8080 8090 8090 8090 GANACICTC ANGRACICTC ACTOCINGLAC CCANCIDATC TICAGCATCT CTTTIQAGA TICCTAGAAT GCGACAACT CTAGGTCAAG CTAGATTGGG TGAGCACGTG GGTTGACTAG AAGTCGTAGAA
7230 TTGGTCATGA			7500 TTATCAGCAA AATAGTCGTT	7590 CGGGAAGCTA GCCCTTCGAT		CCTCCGATCG	7860 GTAAGATGCT CATTCTACGA	7950 ATACGGGATA TATGCCCTAT	8040 CCGCTGTTGA GGCGACAACT
7220 TTAAGGGATT	7310 ATATGAGTAA TATACTCATT	7400 CTGACTCCCC	7490 GGCTCCAGAT CCGAGGTCTA	7570 7580 TCCAGTCTAT TAATTGTTGC AGGTCTAT ATTAACAACG		7760 CTCCTTCGGT		7940 CCCGGCGTCA GGGCCGCAGT	8030 AAGGAICTTA
7210 7220 7220 AAAACTCACG TTAAGGGATT TTGGTCATGA TTTTTTTTTT	7200 TCTAAAGTAT	7390 CCATAGTTGC	7480 CACGCTCACC GTGCGAGTGG	7570 TCCAGTCTAT	7660 TGGTGTCACG	AAGCGGTTAG	7840 CTCTTACTGT	7930 GTTGCTCTTG	BOZO GAAACICTC CTTTTGAGAG

Figure 14I (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement) 8180

CTCA	AAAC	•
ATGTTGAATA TACAACTTAT	8270 TTAGAAAAT AATCTTTTTA	
CGACACGGAA GCTGTGCCTT	8260 TYGAATGTAT AACTTACATA	
GGAATAAGGG	8250 GGATACATAT CCTATGTATA	
CGCAAAAAAG GCGTTTTTTC	8240 TCTCATGAGC AGAGTACTCG	် တ
8140 GGCAAAATGC CCGTTTTACG	AGGTTATTG TCCCAATAAC	GGGTTCCGCC CARARAGTCC CARARAGTCC CACCTCACGT C
8130 AAAACAGGAA TTTTGTCCTT	8220 AGCATTTATC TCGTAAATAG	8310 CGAAAAGTGC
8120 TGGGTGAGCA ACCCACTCGT	8210 ATATTATTGA TATAATAACT	8300 CACATTTCCC
8110 8120 8130 8130 8140 8140 CCAGCGITTION BILL CCAGCGITTIC CONTRAGGG CGACACGGAA ATGTTGAATA CTCA CCAGCGITTIC TGGGTGAGCA ATGTTGATTA GAGTGGCGAAAA ATGTTGTCCTT TACAACTTAT GAGTGGCAAAG ACCCACTCGT TTTTGTCCTT CCGTTTTTTTTTT	8200 8210 8270 8270 8270 8270 8270 7CCTTTTTC AGGGTTATTG TCTCATGAGC GGATACATAT TTGAATGTAT TTAGAAAAAT AAACACTTTTTCA ATATTATTG TCCTTTTTCA CTAAAAAT TCCTTTTTTA AGGAAAAAGT TATAATAACT TCGTAAATAG TCCCAATAAC AGAGTACTTCG CCTATGTATA AACTTACATA AATCTTTTTA TTTCAAAAAAGT TATAATAACT TCGTAAATAG TCCCAATAAC	8290 GGGTTCCGCG

Figure 14J (SEQ ID NO.: 10 – Primary Sequence) (SEQ ID NO.: 28 – Complement)

(SEQ ID NO.: 23 – Primary Sequence) (SEQ ID NO.: 29 – Complement)

550 GGTCTTCTGG CCAGAAGACC	490 AGGCCCCGTC TCCGGGGCAG	430 GCTCAGCGCT CGAGTCGCGA	370 GTGGACAAGA CACCTGTTCT	310 AGCAGCTTGG TCGTCGAACC	250 CCGGCTGTCC GGCCGACAGG	190 CCCGAACCGG GGGCTTGGCC	130 TCCTCCAAGA AGGAGGTTCT	70 TTGGAATTICT AACCTTAAGA	10 GGTACCAATT C
0 G CTTTTTCCCC C GAAAAAGGGG	500 TGCCTCTTCA ACGGAGAAGT	440 CCTGCCTGGA GGACGGACCT	380 AAGTTGGTGA TTCAACCACT	320 GCACCCAGAC CGTGGGTCTG	260 TACAGTCCTC ATGTCAGGAG	200 TGACGGTGTC ACTGCCACAG	140 GCACCTCTGG CGTGGAGACC	BO TGCGGCCGCT '	20 TAAATTGATA 7 ATTTAACTAT <i>1</i>
o 570 C AGGCTCTGGG G TCCGAGACCC	T GGGCCTCCGG	450 CGCATCCCGG CGCTAGGGCC	390 GAGGCCAGCA CTCCGGTCGT	330 CTACATCTGC GATGTAGACG	270 AGGACTCTAC TCCTGAGATG	210 GTGGAACTCA CACCTTGAGT	150 GGGCACAGCG (CCCGTGTCGC (90 TGCTAGCACC 1 ACGATCGTGG 1	30 TCTCCTTAGG T AGAGGAATCC A
580 CAGGCACAGG GTCCGTGTCC	520 TCTGCCCGCC AGACGGGCGG	460 CTATGCAGCC GATACGTCGG	400 CAGGGAGGGA GTCCCTCCCT	340 AACGTGAATC TTGCACTTAG	280 TCCCTCAGCA AGGGAGTCGT	220 GGCGCCCTGA CCGCGGGACT	160 GCCCTGGGCT (CGGGACCCGA	100 AAGGGCCCAT (TTCCCGGGTA (40 TCTCGAGTCT (AGAGCTCAGA (
C GATCCACGGG	CCACTO		GGGTGTCTGC CCCACAGACG	350 ACAAGCCCAG TGTTCGGGTC	290 GCGTGGTCAC CGCACCAGTG	230 CCAGCGGCGT GGTCGCCGCA	170 GCCTGGTCAA CGGACCAGTT	110 CGGTCTTCCC (GCCAGAAGGG	50 CTAGATAACC G GATCTATTGG C
CTAACCCAGG GATTGGGTCC	TCAGGGAC AGTCCCTC	GCAGCAAC	TGGAAGCCAG ACCTTCGGTC	360 CAACACCAAG GTTGTGGTTC	GCACGGGAGG	240 GCACACCTTC CGTGTGGAAG	180 GGACTACTTC CCTGATGAAG	120 CCTGGCACCC GGACCGTGGG	60 GGTCAATCGA CCAGTTAGCT

1150 GTGTGGTCAG CACACCACTIC	1090 . GCGTGGAGGT CGCACCTCCA	1030 GCGTGGTGGT CGCACCACCA	970 TCCTCTTCCC AGGAGAAGGG	910 CTGACACGTC GACTGTGCAG	850 CTCANGGCGG GAGTTCCGCC	790 TGACAAAACT ACTGTTTTGA	730 CCTTCTCTCC GGAAGAGAGG	670 CCCTGCCCCT GGGACGGGGA	610 CCCTGCACAC GGGACGTGTG	
CGTCCT	GCATAA' CGTATT	GGACGT CCTGCA	CCCAAAA GGGTTTT	CACCTCC GTGGAGG	850 CGG GACAGGTGCC GCC CTGTCCACGG	90 PT CACACATGCC 3A GTGTGTACGG	740 C TCCCAGATTC G AGGGTCTAAG	0 T GACCTAAGCC A CTGGATTCGG	620 C AAAGGGGCAG G TYTTCCCCGTC	
GTCCTG	1110 TGCC AAGACAAAGC ACGG TTCTGTTTCG	1040 GAGC CACGAAGACC CTCG GTGCTTCTGG	980 CCC AAGGACACCC GGG TTCCTGTGGG	920 ATC TCTTCCTCAG TAG AGAAGGAGTC	870 CTAGAGTAGC GGATCTCATCG	810 CACCGTGCCC G GTGGCACGGG	0 750 C CAGTAACTCC G GTCATTGAGG	690 CACCCCAAAG G GTGGGGTTTC	630 GIGCTGGGCT CACGACCCGA	
1170 CACC AGGACTGGCT GTGG TCCTGACCGA	10 3C CGCGGGAGGA CG GCGCCCTCCT	1060 CTGAGGTCAA G GACTCCAGTT	1000 C TCATGATCTC G AGTACTAGAG	0 940 G CACCTGAACT C GTGGACTTGA	980 CTGCATCCAG GACGTAGGTC	AGGTAAGCCA TCCATTCGGT	760 CAATCTTCTC GTTAGAAGAG	700 GCCAAACTCT CGGTTTGAGA	640 CAGACCTGCC GTCTGGACGG	
GAATIGG CTTACC	GCAGT: CGTCA	GTTCAA(CAAGTT(CCGGAC		GGAÇAGGO	ecccaggccr cgggrccga	770 TCTGCAGAGC AGACGTCTCG	710 CCACTCCCTC AGCTCGGACA GGTGAGGGAG TCGAGCCTGT	AAGAGCCATA TTCTCGGTAT	
 GAGUAC CICATG		TACGTGC	GAGGTC! CTCCAG	ccercae eecaere	CAGCCGGGTG GTCGGCCCAC	CGCCCTCCAG GCGGGAGGTC	780 CCAAATCTTG GGTTTAGAAC	720 AGCTCGGACA TCGAGCCTGT	CCGGGAC	

1750 ACGTACCCCC TGTACATACT TGCATGGGGG ACATGTATGA	1690 1700 GTGCGACGGC CGGCNAGCCC CACGCTGCCG GCCGTTCGGG	1630 1640 CATGAGGCTC TGCACAACCA GTACTCCGAG ACGTGTTGGT	1570 1580 AAGCTCACCG TGGACAAGAG TTCGAGTGGC ACCTGTTCTC	1520 AACAACTACA AGACCACGCC TTGTTGAIIGT TCTGGTGCGG	1450 GTCAAAGGCT TCTATCCCAG CAGTTTCCGA AGATAGGGTC	1390 1400 TACACCCTGC CCCCATCCCG ATGTGGGACG GGGGTAGGGC	1330 1340 CTGAGAGTGA CCGCTGTACC GACTCTCACT GGCGACATGG	1270 1280 GTGGGACCCG TGGGGTGCGA CACCCTGGGC ACCCCACGCT	312 1210 GCAAGGTCTC CAACAAA CGTTCCAGAG GTTGTTT	(SEQ ID NO.: 29 – Complement)	Figure 19C Figure 19C Figure 19C
1760 ATACT TATGA	1700 AGCCC TCGGG	1640 VACCA	1580 AGAG	1520 CGCC GCGG	1460 CCAG GGTC	1400 CCCG GGGC	1340 PACC ATGG	1280 GCGA CGCT	1220 AGCC TCGG	nent)	quenc
1770 T TCCCGGGCGC A AGGGCCCGCG	0 1710 C CCGCTCCCCG G GGCGAGGGGC	1650 A CTACACGCAG F GATGTGCGTC	1590 CAGGTGGCAG GTCCACCGTC	1530 TCCCGTGCTG AGGGCACGAC	CGACATCGCC GCTGTAGCGG	1410 GGATGAGCTG CCTACTCGAC	1350 AACCTCTGTC TGGAGACAG	1290 GGGCCACATG CCCGGTGTAC	1230 <u>33</u> CAACAAAGCC CTCCCAGCCC C GTTGTTTCGG GAGGGTCGGG G		e)
1780 CCAGCATGGA GGTCGTACCT	1720 GGCTCTCGCG CCGAGAGCGC	1660 AAGAGCCTCT TTCTCGGAGA	1600 CAGGGGAACG GTCCCCTTGC	1540 GACTCCGACG CTGAGGCTGC	1480 GTGGAGTGGG CACCTCACCC	1420 ACCAAGAACC TGGTTCTTGG	1360 CCTACAGGGC GGATGTCCCG	1300 GACAGAGGCC CTGTCTCCGG	1240 CATCGAGAA GITAGCTCTT		
-		1670 CCCTGTCTCC GGGACAGAGG	1610 TCTTCTCATG CTCCGTGATG AGAAGAGTAC GAGGCACTAC	1550 14 1560 GCTCCTTCTT CCTCTACAGC CGAGGAAGAA GGAGATGTCG	1496 1500 AGAGCAATGG GCAGCCGGAG TCTCGTTACC CTCGCCCTC	1440 AGGTCAGCCTI GACCTGCCTG TCCAGTCGGA CTGGACGGAC	1380 AGCCCCGAGA ACCACAGGTG TCGGGGCTCT TGGTGTCCAC	1310 GGCTCGGCCC / CCGAGCCGGG	1250 AACCATCTCC AAAGCCAAAG TTGGTAGAGG TTTCGGTTTC	 pD17-hG1b	-
1790 AATAAAGCAC CCAGCGCTGC TTATTTCGTG GGTCGCGACG	1730 1740 GTCGCACGAG: GATGCTTGGC CAGCGTGCTC CTACGAACCG	1680 GGGTAAATGA CCCATTTACT	1620 CTCCGTGATG GAGGCACTAC	1560 CCTCTACAGC GGAGATGTCG	1500 GCAGCCGGAG CGTCGGCCTC	TA40 GACCTGCCTG CTGGACGGAC	1380 GAGA BACCACAGGTG CTCT TGGTGTCCAC	1320 ACCCTCTGCC TGGGAGACGG	1260 PAAAGCCAAAG BATTTCGGTTTC	 	

Figure 19D (SEQ ID NO.: 23 – Primary Sequence) (SEQ ID NO.: 29 – Complement)

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2350	2290	2230	2170	2110	2050	1990	1930	1870	1810
GCTTTCCCCG	CCAGCGCCCT	GTAGCGGCGC	CTATGGCTTC	GCGCCCCTGT	GGAGCCCCTG	TTGCCAGCGT	TGTGCCTGGG	AGTGGCATGA	CCTGGGCCCC
CGAAACGGGC	GGTCGCGGGA	CATCGCCGCG	GATACCGAAG	CGCGGGGACA	CCTCGGGGAC	AACGGTCGCA	ACACGGACCC	TCACCGTACT	GGACCCGGGG
2360	2300	2240	2180	2120	2060	2000	1940	1880	1820
TCAAGCTCTA	AGCGCCCGCT	ATTAAGCGCG	TGAGGCGGAA	CCTCCCGACC	GGGACAGACA	GGCCCTCCCT	CCCCCTAGGG	GGGAGGCAGA	TGCGAGAC'FG
AGTTCGAGAT	TCGCGGGCGA	TAATTCGCGC	ACTCCGCCTT	GGAGGGCTGG	CCCTGTCTGT	CCGGGAGGGA	GGGGGATCCC	CCCTCCGTCT	ACGCTCTGAC
2370	2310	2250	2190	2130	2070	2010	1950	1890	1830
AATCGGGGCA	CCTTTCGCTT	GCGGGTGTGG	AGAACCAGCT	TCCATGCCCA	CACAGCCCCT	CCAGCAGCAC	TGGGGCTCAG	GCGGGTCCCA	TGATGGTTCT
TTAGCCCCGT	GGAAAGCGAA	CGCCCACACC	TCTTGGTCGA	AGGTACGGGT	GTGTCGGGGA	GGTCGTCGTG	ACCCCGAGTC	CGCCCAGGGT	ACTACCAAGA
2380	2320	2260	2200	2140	2080	2020	1960	1900	1840
TCCCTTTAGG	TCTTCCCTTC	TGGTTACGCG	GGGGCTCTAG	CTCGGGGGCA	GCCTCTGTAG	CTGCCCTGGG	CCAGGGGCTG	CTGTCCCCAC	TTCCACGGGT
AGGGAAATCC	AGAAGGGAAG	ACCAATGCGC	CCCCGAGATC	GAGCCCCCGT	CGGAGACATC	GACGGGACCC	GGTCCCCGAC	GACAGGGGTG	AAGGTGCCCA
2390	2330	2270	2210	2150	2090	2030	1970	1910	1850
CGATTT	CTTTCTCGCC	CAGCGTGAGG	GGGGTATCCC	TGCTGGGGAT	GAGAÇTGTÇÇ	CTGGGCCACG	CCCTCGGCAG	ACTGGCCCAG	CAGGĆCGAGT
CCTAAA	GAAAGAGCGG	GTCGCACTGG	CCCCATAGGG	ACGACCCCTA	CTCTGACAGG	GACCCGGTGC	GGGAGCCGTC	TGACCGGGTC	GTCCGGCTCA
2400	2340	2280	2220	2160	2100	2040	1980	1920	1860
AGTGCTTTAC	ACGTTCGCCG	GCTACACTTG	CACGCGCCCT	GCGGTGGGCT	TGTTCTGTGA	GGAAGCCCTA	GGTGGGGGAT	GCTGTGCAGG	CTGAGGCCTG
TCACGAAATG	TGCAAGCGGC	CGATGTGAAC	GUGCGCGGGA	CGCCACCCGA	ACAAGACACT	CCTTCGGGAT	CCACCCCCTA	CGACACGTCC	GACTCCGGAC
					;		•		•

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2950 CGCGCCAAAC GCGCGGTTTTG	2890 GAGGCTTTTTT CTCCGAAAAA	2830 Typyyyatyya Aaaaataaat	2770 CCATCCCGCC GGTAGGGCGG	2710 GAAGTATGCA CTTCATACGT	2650 AATTCTGTGG TTAAGACACC	2590 TGGGGATTTC ACCCCTAMAG	2530 TYCCAAAC'IGG AGGTTTTGACC	2470 GATAGACGGT ' CTATCTGCCA	2410 GGCACCTCGA (CCGTGGAGCT ((SEQ ID NO.: 23 – Primary Sequence) (SEQ ID NO.: 29 – Complement)
2960 TTGACGGCAA	2900 TGGAGGCCTA ACCTCCGGAT	2840 TGCAGAGGCC ACGTCTCCGG	2780 CCTAACTCCG GGATTGAGGC	2720 AAGCATGCAT TTCGTACGTA	2660 AATGTGTGTC TTACACACAG	2600 GGCCTATTGG CCGGATAACC	2540 AACAACACTC TTGTTGTGAG	2480 TTTTCGCCCT ' AAAAGCGGGA	2420 CCCCAAAAAA (GGGGTTTTTTT (rigure 195 : 23 – Primary Sequer O.: 29 – Complement
2970 TCCTAGCGTG AGGATCGCAC	2910 GGCTTTTGCA CCGAAAACGT	2850 GAGGCCGCCT CTCCGGCGGA	2790 CCCAGTTCCG GGGTCAAGGC	2730 CTCAATTAGT GAGTTAATCA	2670 AGTTAGGGTG TCAATCCCAC	2610 TTAAAAATG AATTTTTTAC	2550 AACCCTATCT (TIGGGATAGA (2490 TTGACGTTGG 1 AACTGCAACC !	2430 CTTGATTAGG (GAACTAATCC (nce)
2980 AAGGCTGGTA TTCCGACCAT	2920 AAAAGCTTGG TTTTCGAACC	2860 CGGCCTCTGA GCCGGAGACT	2800 CCCATTCTCC GGGTAAGAGG	2740 CAGCAACCAT GTCGTTGGTA	2680 TGGAAAGTCC ACCTTTCAGG	2620 AGCTGATTTA TCGACTAAAT	2560 CGGTCTATTC GCCAGATAAG	2500 AGTCCACGTT (TCAGGTGCAA (2440 GTGATGGTTC I	
 2990 GGATTTTATC CCTAAAATAG	2930 ACAGCTCAGG TGTCGAGTCC	2870. 2870. CEATAGTAGTAGTAGCAGCCGATAAGGTC TTCATCACTC	2000 2000 2010	2750 AGTCCCGCCC TCAGGGCGGG	2690 CCAGGCTCCC GGTCCGAGGG	2630 ACAAAAATTT TGTTTTTAAA	2570 TTTTGATTTA (AAAACTAAAT)	2510 [†] CTTTAATAGT (GAAATTATCA (2450 ACGTAGTEGG CCATCGCCCT TGCATCACCC GGTAGCGGGA	pD17-hG1b
3000 CCCCCTCCCA CCCCCCCCTCCCCTCCCCCCCCCCC	2940 ACAGCTCAGG GCTGCGATTT TGTCGAGTGG CGACGCTAAA	2880 AAGTAGTGAG TTCATCACTC	2820 TGACTAATTT ACTGATTAAA	2760 CTAACTCCGC GATTGAGGCG	2700 CAGGCAGGCA GTCCGTCCGT	2640 AACGCGAATT TTGCGCTTAA	2580 TAAGGGATTT ATTCCCTAAA	2520 GGACTCTTGT CCTGAGAACA	2460 CATCGCCCT GTAGCGGGA	

3550 TTGAAGTCTA NACTTCAGNT	3490 TCCCAGAATA AGGGTCTTAT	3430 TGCAGGAATT' ACGTCCTTAA	3370 CTGTTTACCA GACAAATGGT	3310 TTATTGAACA AATAACTTGT	3250 TCAAAGAACC AGTTTCTTGG	3190 CCATTCCTGA GGTAAGGACT	3130 CAACCTCTTIC GTTGGAGAAG	3070 ACGGAGACCT TGCCTCTGGA	JO10 TCATGGTTCG AGTACCAAGC	
0 A CGAGAAGAAA I' GCTCTTCTTT	3500 A CCCAGGCGTC T GGGTCCGCAG	3440 TGAAAGTGAC ACTTTCACTG	3380 GGAAGCCATG CCTTCGGTAC	3320 ACCGGAATTG TGGCCTTAAC	3260 ACCACGAGGA TGGTGCTCCT	3200 GAAGAATCGA CTTCTTAGCT	3140 AGTGGAAGGT TCACCTTCCA	3080 ACCCTGGCCT TGGGACCGGA	3020 ACCATTGAAC '	
3570 A GACTAACAGG T CTGATTGTCC	3510 CTCTCTGAGG GAGAGACTCC	3450 ACGTTTTTCC TGCAAAAAGG	3390 AATCAACCAG TTAGTTGGTC	3330 GCAAGTAAAG CGTTCATTTC	3270 GCTCATTTTC CGAGTAAAAG	3210 CCTTTAAAGG GGAAATTTCC	3150 AAACAGAATC ' TTTGTCTTAG	3090 CCGCTCAGGA 1 GGCGAGTCCT 1	3030 TGCATCGTCG (ACGTAGCAGC (
3580 AAGATGCTTT TTCTACGAAA	3520 TCCAGGAGGA AGGTCCTCCT	3460 CAGAAATTGA GTCTTTAACT	3400 GCCACCTTAG CGGTGGAATC	3340 TAGACATGGT ATCTGTACCA	3280 TTGCCAAAAG AACGGTTTTC	3220 ACAGAATTAA TGTCTTAATT	3160 TGGTGATTAT A	3100 ACGAGTTCAA (TGCTCAAGTT (3040 CCGTGTCCCA <i>I</i> GGCACAGGGT '	
3590 CAAGTTCTCT GTTCAAGAGA		3470 TTTGGGGAAA AAACCCCTTT	3410 ACTCTTTGTG TGAGAAACAC	3350 TTGGATAGTC AACCTATCAG	3290 TTTGGATGAT GCCTTAAGAC AAACCTACTA CGGAATTCTG	3230 TATAGTTCTC AGTAGAGAAC ATATCAAGAG TCATCTCTTG		GTACTTCCAA, CATGAAGGTT,	3050 AAATATGGGG / TTTATACCCG	
GCTCCCCTCC CGAGGGGAGG	AAGTATI TTCATA	TATAAA(ATATTTY	3420 ACTCTTTGTG; ACAAGGATCA TGAGAAACAC TGTTCCTAGT	3360 GGAGGCAGTT CCTCCGTCAA	3300 GCCTTAAGAC CGGAATTCTG	3240 AGTAGAGAAC TCATCTCTTG	3180 ACCTGGTTCT TGGACCAAGA	3120 AGAATGACCA TCTTACTGGT	3060 ATTGGCAAGA TAACCGTTCT	•

Figure 19F (SEQ ID NO.: 23 – Primary Sequence) (SEQ ID NO.: 29 – Complement) AGGTGTGTCC TCCACACAGG

CATAGAGTGT CTGCTATTAA TAACTATGCT GTATCTCACA GACGATAATT ATTGATACGA

CTGCTATTAA TAACTATGCT

4170

4180

CAAAATTGT' GTACCTTTAG

4190

GITTTTAACA : CATGGAAATC

4160

TTCTGTAACC AAGACATTGG

TTTATAAGTA AAATATTCAT

GGCATAACAG CCGTATTGTC

TTATAATCAT AACATACTGT TTTTTCTTAC AATATTAGTA TTGTATGACA ; AAAAAAGAATG

4110

4120

4130

4140

ACGA'TAAATG TGCTATTTAC

TGGTGTTTCC

(SEQ ID NO.: 23 - Primary Sequence) (SEQ ID NO.: 29 - Complement) Figure 19G

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CATTTTTATA	362Ő		
AGACCATGGG	3630		
CATTITITATIA AGACCATIGGG ACTITITIGCTIG GCTTTAGATO, TCTTTIGTGAA	3640		
GCTTTA		٠.	
SATC TO	3650	•	-
TTTGTGAA	3660		

TAAGGTAAAT

ATAAAATTTT

TATTTAAAA ATTCACATAT TACACAATTT

TAAGTGTATA

ATGTGTTAAA

GATGACTAAG ATTAACAAAC

CTACTGATTC TAATTGTTTG

3730

3740

3750

3760

3770

3780

ATTCCATTTA

ACATAAAATC

TGTATTTAG

3810 3820 ATTCCAACCT ATGGAACTGA TGAATGGGAG TAAGGTTGGA TACCTTGACT ACTTACCCTC

CAGTGGTGGA ATGCCTTTAA

3830

3840

3790

TGAGGAAAAC

CTGTTTTGCT GACAAAAACGA

GTCTTCTTTA CGGTAGATCA

CAGAAGAAAT

GCCATCTAGT

GATGÄTGAGG

CTACTGCTGA

3900

CTACTACTCC, GATGACGACT

3850

3860

3870

3880

3890

ACTCCTTTTG

CTCTCAACAT TCTACTCCTC
GAGAGTTGTA AGATGAGGAG

GITTTTTCTT

CTCTTTCCAT

CAAAAAAGAA GAGAAAGGTA

GAAGACCCCA AGGACTTTCC

CTTCTGGGGT TCCTGAAAGG

3910

3920

3930

3940

3950

3960

NAGTCTTYAAC TTCAGAATTG

GATTCAAAAA CTAAGTTTTT

TGAGTCATGC ACTCAGTACG

ACACAAATCA TGTGTTTAGT

AATAGAACTC TTGCTTGCTT

4030

ACCACAAAGG

AAAAAGCTGC TTTTTTCGACG

ACTGCTATAC TGACGATATG

AAGAAATTA TGGAAAAATA

TTCTTTTAAT ACCTTTTTAT

4060

4070

4080

3970

3980

3990

4000

4010

4020

CCTTGGAATG GGAACCTTAC

AAGACACCAC

ACTGTATTAA

TGACATAATT

GGACAAACTA

CCTGTTTGAT

GGATGTCTCT

AAATTTCGAG

CCTACAGAGA TTTAAAGCTC

TTCTGTGGTG

ATTTCGATAC TAAAGCTATG

GTAAAAATAT TCTGGTACCC

TGAAAACGAC ACTTTTGCTG

CGAAATCTAG 'AGAAACACTT GCTTTAGATC, TCTTTGTGAA

3670

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3690

3700

3710

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Figure 19H
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

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4750 TGTGAAATTG ACACTTTAAC	TCTGTATACC AGACATATGG	4630 CΑΊΨΤΙΊΤΤΤΟ GΊΡΑΑΑΑΛΑΑG	4570 TTATTGCAGC AATAACGTCG	4510 GCTGGATGAT CGACCTACTA	4450 CACTGCATTC GTGACGTAAG	4390 CTTATAATGG GAATATTACC	4330 TCCCCCTGAA AGGGGGACTT	4270 TCATAATCAG AGTATTAGTC	4210 CTTTTTAATT GAAAAATTAA
4760	4700	4640	4580	4520	4460	4400	4340	4280	4220
TTATCCGCTC	GTCGACCTCT	ACTGCATTCT	TTATAATGGT	CCTCCAGCGC	TAGTTGTGGT	TTACAAATAA	CCTGAAACAT	CCATACCACA	TGTAAAGGGG
AATAGGCGAG	CAGCTGGAGA	TGACGTAAGA	AATATTACCA	GGAGGTCGCG	ATCAACACCA	AATGTTTATT	GGACTTTGTA	GGTATGGTGT	ACATTTCCCC
4770 ACAATTCCAC TGTTAAGGTG	AGCTAGAGCT TCGATCTCGA	4650 AGTTGTTGGTT TCAACACCAA	4590 ТАСАРАТАРА АТСТТТАТТТ	4530 GGGGATCTCA CCCCTAGAGT	4470 TTGTCCAAAC AACAGGTTTG	4410 AGCAATAGCA TCGTTATCGT	4350 AAAATGAATG ITTTACTTAC	4290 TTTGTAGAGG AAACATCTCC	4230 TTAATAAGGA AATTATTCCT
4780	4720	4660	4600	4540	4480	4420	4360	4300	4240
ACAACATACG	TGGCGTAATC	TGTCCAAACT	GCAATAGCAT	TGCTGGAGTT	TCATCAATGT	TCACAAATTT	CAATTGTTGT	TTTTACTTGC	ATATTTGATG
TGTTGTATGC	ACCGCATTAG	ACAGGTTTGA	CGTTATCGTA	ACGACCTCAA	AGTAGTTACA	AGTGTTTAAA	GTTAACAACA	AAAATGAACG	TATAAACTAC
4800	4730	4670	4610	4550	4490°; 4500	4440	4380	4310	4250
AGCCGGAAGC ATAAAGTGTA	ATGGTCATAG	CATCAATGTA	CACAAATTTC	CTTCGCCCAC CCCAACTTGT	АТСТТАТСАТ, СТСТСВСАТСС	САСАААТААА GCATTTTTTT	TGTTAACTTG TTTATTGCAG	TITAAAAAAC	TATAGTGCCT
TCGGCCTTCG TATTTCACAT	TACCAGTATC	GTAGTTACAT	GTGTTTAAAG	GAAGCGGGTG GGGTTGAACA	ТАСААТАСТА, САСАССТАСС	GTGTTTATTT (CGTAAAAAAA	ACAATTGAAC NANTAACGTC	AAATTTTTTG	ATATCACGGA
4800	4740	4680	4620	4560	4500	4440	4380	4320	4260
ATAAAGTGTA	CTGTTTCCTG	TGTTATCATG	ACAAATAAAG	CCCAACTTGT	GTCTGGATCG	GCATTTTTTT	TYTTATYGCAG	CTCCCACACC	TGACTAGAGA
TAVITITCACAT	GACAAAGGAC	AGAATAGTAC	TGTTTATTTC	GGGTTGAACA	CAGACCTAGC	CGTAAAAAAA	NANTAACGTC	GAGGGTGTGG	ACTGATCTCT

(SEQ ID NO.: 23 - Primary Sequence) (SEQ ID NO.: 29 - Complement)	HIGHTP 191
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	Q ID NO.: 29 – Complement)	ID NO.: 23 - Primary Sequence)
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			4840 TCACATTIAAT AGTGTAATTA 4900 TGCATTAATTAATG ACGTAATTAC 4960 CTTCCTCGCT GAAGGAGCAAAAGG CTCGAAAAGG CTCGTTTCCG TGAGCTAAAAGG CTCGTTTCCG 5080 GAGCAAAAGG CTCGTTTCC 5140 ATAGGCTCCGAGGC TATCCGAGGCC	4850 'IGCGTTIGCGC TIC ACGCAACGC ACGCAACGC AATTCGGCCAACCCTTAAGCCGGTT GC GTGACTGACTGAGC GC GGTAATACGG TY CCATTATACGG TY CCAGCAAAAG GGGGTCGTTTTC CC GGGGGGACTG CC GGGGGGACTG CC GGGGGGACTG CC GGGGGGACTG CC ACTATAAAGA T	4860 TGACTGCCCG AGJIGACGGCC AGJIGACGGGA GCGCGCGCCCCT 4980 CTGCGCGCCCCCT 5040 TTATCCACAG AATAGGTGTC CGGTCCTTGG GACCATCACA CTCGTAGTGTCT 5220 TACCAGGCGT
5110	5120	5130.	5140		5160
STAAAAAGGC	CGCGTTGCTG	GCGTTTTTTCC	ATAGGCTCCG		GAGCATCACA
CATTTTTCCCG	GCGCAACGAC	CGCAAAAAAGG	TATCCGAGGC		CTCGTAGTGT
5170	5180	5190	5200	5210	5220
AAATCGACG	CTCAAGTCAG	AGGTGGCGAA	ACCCGACAGG	NAAGA	TACCAGGCGT
TTTAGCTGC	GAGTTCAGTC	TCCACCGCTT	TGGGCTGTCC	TTTCT	ATGGTCCGCA
5230	5240	5250	5260	5280	5280
TTCCCCCTGG	AAGCTCCCTC	GTGCGCTCTC	CTGTTCCGAC	CCTGCCGCTT.JACCGGATACC	ACCGGATACC
AAGGGGGACC	TTCGAGGGAG	CACGCGAGAG	GACAAGGCTG	GGACGCGAA. TGGCCTATGG	TGGCCTATGG
5290	5300	5310	5320	5330	5340
TGTCCGCCTT	TCTCCCTTCG	GGAAGCGTGG	CGCTTTCTCA	ATGCTCACGC TCTAGGTATC	TGTAGGTATC
ACAGGCGGAA	AGAGGGAAGC	CCTTCGCACC	GCGAAAGAGT	TACGAGTGCG ACATCCATAG	ACATCCATAG
5350 TCAGTTCGGT	5360 GTAGGTCGTT CATCCAGCAA	5370 CGCTCCAAGC GCGAGGTTCG	5380 TGGGCTGTGT ACCCGACACA	GCACGAACCC CGTGCTTGGG	GCCGTTCAGC

5990 5900 5910 5920 5930 5940 ACAGTYACCA ATGCTTAATC AGTGAGGCAC CTATCTCAGC GATCTGTCTA TTTCGTTCAT TGTCAATGGT TACGAATTAG TCACTCCGTG GATAGAGTCG CTAGACAGAT AAAGCAAGTA

5950 5960 5970 5980 5990 5990 6000 CCATAGTIGG CTGACTCCC GTCGTGTAGA TAACTACGAT ACGGGAGGGC TTACCATCTG GGTAGTCT ATTGATGCTA TGCCCTCCCG AAIGGTAGAC

Figure 19J
(SEQ ID NO.: 23 – Primary Sequence)
(SEQ ID NO.: 29 – Complement)

	5830 Titraaatita Aaatittaat	5770 NAANCTCACG TYTTGAGTGC	5710 AAAAAGGATC TTTTTTTCCTAG	5650 AACAAACCAC TTGTTTGGTG	5590 · TCTGCGCTCT AGACGCGAGA	5530 CTACAGAGTT GATGTCTCAA	5470 TATCGCCACT (ATAGCGGTGA (5410 CCGACCGCTG (GGCTGGCGAC (•	(SEQ ID NO.: 29 - Complement)	SEO ID NO.: 23 - Primary Sequence)
	5840 AAAATGAAGT TTTTACTTCA	5780 TTAAGGGATT AATTCCCTAA	5720 TCAAGAAGAT AGTTCTTCTA	5660 CGCTGGTAGC GCGACCATCG	5600 GCTGAAGCCA CGACTTCGGT	5540 CTTGAAGTGG GAACTTCACC	5480 GGCAGCAGCC CCGTCGTCGG	5420 CGCCTTATCC GCGGAATAGG		- Complement	Primary Sequen
	5850 TTTAAATCAA AAATTTAGTT	5790 TTGGTCATGA AACCAGTACT	5730 CCTTTGATCT GGAAACTAGA	5670 GGTGGTTTTT CCACCAAAAA	5610 GTTACCTTCG CAATGGAAGC	5550 TGGCCTAACT ACCGGATTGA	5490 ACTGGTAACA TGACCATTGT	5430 GGTAACTATC CCATTGATAG		,	ice)
	5060 TCTAAAGTAT AGATTTCATA	5800 GATTATCAAA CTAATAGTTT	5740 TTTCTACGGG AAAGATGCCC	5680 TTGTTTGCAA AACAAACGTT	5620 GAAAAAGAGT CTTTTTCTCA	5560 ACGGCTACAC TGCCGATGTG	5500 GGATTAGCAG CCTAATCGTC	5440 GTCTTGAGTC CAGAACTCAG			
•.	5870 ATATGAGTAA TATACTCATT		STETERCECT CAGACTECA	5700 GCAGCAGATT! ACGCGCAGAA CGTCGTCTAA: TGCGCGTCTT	5640 TGGTAGCTCTUTGATCCGGCA ACCATCGAGA ACTAGGCCGT	5570 5580 TAGAAGGACA GTATTTGGTA ATCTTCCTGT CATAAACCAT	5510 TO THE STATE OF THE STATE	5450 CAACCCGGTA, AGACACGACT GTTGGGCCAT TCTGTGCTGA	· • •	pD17-hG.1b	
	5880 ACTTGGTCTG TGAACCAGAC	5810 AAGGATCTTC ACCTAGATCC TTCCTAGAAG TGGATCTAGG	5760 CAGTGGAACG GTCACCTTGC	5700 ACGCGCAGAA TGCGCGTCTT	5640 TGATCCGGCA ACTAGGCCGT	5580 GTATTTGGTA CATAAACCAT	# 5520 GTAGGCGGTG CATCCGCCAC	5460 AGACACGACT TCTGTGCTGA	<u> </u>		<u> </u>

Figure 19K (SEQ ID NO.: 23 – Primary Sequence) (SEQ ID NO.: 29 – Complement)

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6550	6490	6430	6370	6310	6250	6190	6130	6070	6010
TGCTCATCAT	GTTGCTCTTG	ТТТСТСТСТСАС	CACTCATGGT	AAGCGGTTAG	CATTCAGCTC	GCAACGTTGT	TCCAGTCTAT	TAAACCAGCC	GCCCCAGTGC
ACGAGTACTA	CAACGAGAAC	АЛАСАСАСТС	GTGAGTACCA	TTCGCCAATC	GTAAGTCGAG	CGTTGCAACA	AGGTCAGATA	ATTTGGTCGG	CGGGGTCACG
6560	6500	6440	6380	6320	6260	6200	6140	6080	6020
TGGAAAACGT	CCCGGCGTCA	TGGTGAGTAC	TATGGCAGCA	CTCCTTCGGT	CGGTTCCCAA	TGCCATTGCT	TAATTGTTGC	AGCCGGAAGG	TGCAATGATA
ACCTTTIGCA	GGGCCGCAGT	ACCACTCATG	ATACCGTCGT	GAGGAAGCCA	GCCAAGGGTT	ACGGTAACGA	ATTAACAACG	TCGGCCTTCC	ACGTTACTAT
6570 TCTTCGGGGC AGAAGCCCCG	6510	6450	6390	6330	6270	6210	6150	6090	6030
	ATACGGGATA	TCAACCAAGT	CTGCATAATT	CCTCCGATCG	CGATCAAGGC	ACAGGCATCG	CGGGAAGCTA	GCCGAGCGCA	CCGCGAGACC
	TATGCCCTAT	AGTTGGTTCA	GACGTATTAA	GGAGGCTAGC	GCTAGTTCCG	TGTCCGTAGC	GCCCTTCGAT	CGGCTCGCGT	GGCGCTCTGG
6580	6520	6460	6400	6340	6280	6220	6160	6100	6040
GAAAACTCTC	ATACCGCGCC	CATTCTGAGA	CTCTTACTGT	TTGTCAGAAG	GAGTTACATG	TGGTGTCACG	GAGTAAGTAG	GAAGTGGTCC	CACGCTCACC
CTTTTGAGAG	TATGGCGCGG	GTAAGACTCT	GAGAATGACA	AACAGTCTTC	CTCAATGTAC	ACCACAGTGC	CTCATTCATC	CTTCACCAGG	GTGCGAGTGG
	6530 ACATAGCAGA TGTATCGTCT	ATAGTGTATG TATCACATAC	CATGCCATCC CTACGGTAGG	6350; TAAGTTGGCC ATTCAACCGG	ATCCGCCATC TAGGGGGTAC	CTCGTCGTTT GAGÇAGCAAA	6170 TTCGCCAGTT AAGCGGTCAA	6110. TGCAACTTTA TCCGCCTCCA ACGTTGAAAT AGGCGGAGGT	6050 GGCTCCAGAT CCGAGGTCTA
6590 AAGGATCTTA CCGCTGTTGA TYCCTAGAAT GGCGACAACT	6540 ACTTTAAAAG TGAAATTTTC	CGGCGACCGA GCCGCTGGCT	6420 GTAAGATGCT CATTCTACGA	6360 GCAGTGTTAT CGTCACAATA	6300 TITGTGCAAAA AACACGTTTTT	6240 GGTATGGCTT CCATACCGAA	6180 AATAGTTTGC TTATCAAACG	6120 TCCGCCTCCA AGGCGGAGGT	6060 TTATCAGCAA AATAGTCGTT

Figure 19L (SEQ ID NO.: 23 – Primary Sequence) (SEQ ID NO.: 29 – Complement)

		,							
7150 CAATTGCATG	7090 GGAGGTCGCT CCTCCAGCGA	7030 CAGTACAATC GTCATGTTAG	6970 TTYATTYTAT AAATAAATA	6910 TAGGTGACCT ATCCACTGGA	6850 GGGTTCCGCG CCCAAGGCGC	6790 AGGGTTATTG TCCCAATAAC	6730 CGACACGGAA GCTGTGCCTT	6670 CCAGCGTITTC GGTCGCAAAG	6610 GATCCAGTTC CTAGGTCAAG
0 G AAGAATCTGC C TYCTTAGACG	7100 GAGTAGTGCG	7040 TGCTCTGATG ACGAGACTAC	6980 TTTTGAGATG AAAACTCTAC	6920 GAGGCGCGCC CTCCGCGCGCGG	6860 CACATTTCCC GTGTAAAGGG	6800 TCTCATGAGC AGAGTACTCG	6740 ATGTTGAATA TACAACTTAT	6680 TGGGTGAGCA ACCCACTCGT	6620 GATGTAACCC CTACATTGGG
7170 TTAGGGTTAG	7110 GCTCGTTTTA	7050 CCGCATAGTT GGCGTATCAA	6990 GAGTTTGGCG CTCAAACCGC	6930 GGCTTCGAAT CCGAAGCTTA	6870 CGAAAAGTGC GCTTTTCACG	6810 GGATACATAT CCTATGTATA	6750 CTCATACTCT GAGTATGAGA	6690 AAAACAGGAA TTTTGTCCTT	ACTCGTGCAC TGAGCACGTG
7180 GCGTTTTGCG CGCAAAACGC	7120 TTAAGCTACA AATTCGATGT	7060 AAGCCAGTAT TTCGGTCATA	7000 CCGATCTCCC GGCTAGAGGG	AGCCAGAGTA TCGGTCTCAT	6880 CACCTGACGT GTGGACTGCA	6820 TTGAATGTAT AACTTACATA	6760 TCCTTTTTCA AGGAAAAAGT	6700 GGCAAAATGC CCGTTTTACG	6640 CCAACTGATC GGTTGACTAG
		-	7010 GATCCCCTAT GGT CTAGGGGATA CC	6950. ACCTTTTTTTT PTAATTTTAT TGGAAAAAAA AATTAAAATA	6890 CGACGGATCG GCTGCCTAGC	6830 TTAGAAAAAT AATCTTTTTA	6770 ATATTATTGA TATAATAACT	6710 CGCAAAAAAG GCGTTTTTTTC	TTCAGCATCT (
7200 CTGCTTCGCG) ATGTACGGGC GACGAAGCGC TACATGCCCG	7130 : 7140 ACAAGGCAAG* GCTTGACCGA TGTTCCGTTC CGAACTGGCT	ACAC!	7010 7020 GATCCCCTAT GETCGACTCT CTAGGGGATA CCAGCTGAGA	6960 FTAATTTTAT AATTAAATA	6900 GGAGATCTGC CCTCTAGACG	6840 AAACAAATAG TTTGTTTATC	6780 AGCATTTATC TCGTAAATAG	GGAATAAGGG CCTTATTCCC	6660 TTTACTTTCA AAATGAAAGT

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CCCATTGACG CAAATGGGCG GTAGGCGTGT ACGGTGGGAGGGGTAACTGC GTTTACCCGC CATCCGCACA TGCCACCCTC

Figure 19M (SEQ ID NO.: 23 – Primary Sequence) (SEQ ID NO.: 29 – Complement)

TA	CA CA	
TAGT'I	GATAT CTATA	
7270 TCAT AGTA	7210 TACG ATGC	
AGCCCATATA TCGGGTATAT	7220 CGTTGACATT GCAACTGTAA	
AT 78	ଜନ୍ୟ ଥିଲ ଜନ	
GGAGTTCCG CCTCAAGGC	7230 ATTATTGAC TAATAACTG	
CGTTACATAA GCAATGTATT	7240 TAGTTATTAA ATCAATAATT	* .
7270 7280 7290 7290 7290 7270 7270 7270 7270 727	7210 7220 7230 7240 7250 7260 CAGATATACG CGTTGACATT GATTATTGAC TAGTTATTAA TAGTAATCAA TTACGGGGTC GTCTATATGC GCAACTGTAA CTAATAACTG ATCAATAATT A1CATTAGTT; AATGCCCCAG	- · · ·
55555555555555555555555555555555555555	7260 GGGGTC GCCCAG	

AACGCCAATA

GGGACTTTCC

TAACTGCAGT

ATGGGTGGAC TACCCACCTG

TATTTACGGT AAACTGCCCA ATAAATGCCA TTTGACGGGT

CCCTGAAAGG

7390

7400

7410

7420

7430

7440

TTGCGGTTAT

GAACCGTCAT

GTAGTTCACA

ATCATATGCC TAGTATACGG

AAGTACGCCC TTCATGCGGG

GGATAACTGC AGTTACTGCC

CCTATTGACG

TCAATGACGG

7470

CTTGGCAGTA

7450

7460 CATCAAGTGT

TAAATGGCCC ATTTACCGGG

GCCTGGCATT
CGGACCGTAA

ATGCCCAGTA

TACGGGTCAT GTACTGGAAT ACCCTGAAAG

CATGACCTTA

TGGGACTTTG CTACTTGGCA

GATGAACCGT

7550

7540

GTACATCTAC CATGTAGATG

GTATTAGTCA CATAATCAGT

TCGCTATTAC AGCGATAATG

CATGGTGATG GTACCACTAC

CGGTTTTGGC AGTACATCAA GCCAAAACCG TCATGTAGTT

7580

7590

7600

7610

ACCCGCACC'F AFCGCCAAAC

TGAGTGCCCC

ATTTCCAAGT TAAAGGTTCA

GAGGTGGGGTEAACTGCAGTT

CTCCACCCCA TTGACGTCAA

7670

7680

ACTCACGGG

7650

7660

TGGGAGTTTG ACCCTCAAAC

TTTTGGCACC AAAACCGTGG

TTTTAGTTGC

AAAATCAACG

GGACTTTCCA

CCTGAAAGGT

AAATGTCGTA ACAACTCCGC

CAGATATATA GCAGAGCTCT

77907

7700

7710

7720

7730

TGGGCGTGGA

TAGCGGTTTG

7630

ACCGAC'I'GGC

GCGITGCTGG

GGGCGGGTAA CTGCAGTTAT

TACTGCATAC AAGGGTATCA

CCCAACGACC

CCCGCCCATT

GACGTCAATA

ATGACGTATG

TTCCCATAGT

7370

TGGCTGACCG

Figure 19N (SEQ ID NO.: 23 – Primary Sequence) (SEQ ID NO.: 29 – Complement)

7810 7820 7830 7840 7850 7860 CTGGCTAACT AGAGAACCCA CTGCTTACTG GCTTATCGAA ATTAATACGA CTCACTATAG GACCGATTGA TCTCTTGGGT GACGAATGAC CGAATAGCTT TAATTATGCT GAGTGATATC

7870 GGAGACCCAA GCTT CCTCTGGGTT CGAA

7880

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